

#### Sequence Listing

#### SEQUENCE LISTING

- (1) GENERAL INFORMATION:
  - (i) APPLICANT: Ashkenazi, Avi J.
  - (ii) TITLE OF INVENTION: Apo-2 LI AND Apo-3 POLYPEPTIDES
  - (iii) NUMBER OF SEQUENCES: 28
    - (iv) CORRESPONDENCE ADDRESS:
      - (A) ADDRESSEE: Genentech, Inc.
      - (B) STREET: 1 DNA Way
      - (C) CITY: South San Francisco
      - (D) STATE: California
      - (E) COUNTRY: USA
      - (F) ZIP: 94080
    - (v) COMPUTER READABLE FORM:
      - (A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
      - (B) COMPUTER: IBM PC compatible
      - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
      - (D) SOFTWARE: WinPatin (Genentech)
    - (vi) CURRENT APPLICATION DATA:
      - (A) APPLICATION NUMBER: 09/993234
      - (B) FILING DATE: 19-NOV-01
      - (C) CLASSIFICATION:
  - (vii) PRIOR APPLICATION DATA:
    - (A) APPLICATION NUMBER: 08/828683
    - (B) FILING DATE: 31-MAR-1997
  - (vii) PRIOR APPLICATION DATA:
    - (A) APPLICATION NUMBER: 08/625328
    - (B) FILING DATE: 1-Apr-1996
  - (vii) PRIOR APPLICATION DATA:
    - (A) APPLICATION NUMBER: 08/710802
    - (B) FILING DATE: 23-Sep-1996
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: Marschang, Diane L.
  - (B) REGISTRATION NUMBER: 35,600
  - (C) REFERENCE/DOCKET NUMBER: P1007P1D1
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: 650/225-5416
    - (B) TELEFAX: 650/952-9881
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 181 amino acids
    - (B) TYPE: Amino Acid
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Met Glu Gln Arg Pro Arg Gly Cys Ala Ala Val Ala Ala Ala Leu Leu Leu Val Leu Leu Gly Ala Arg Ala Gln Gly Gly Thr Arg Ser Pro Arg Cys Asp Cys Ala Gly Asp Phe His Lys Lys Ile Gly Leu Phe Cys Cys Arg Gly Cys Pro Ala Gly His Tyr Leu Lys Ala Pro Cys Thr Glu Pro Cys Gly Asn Ser Thr Cys Leu Val Cys Pro Gln Asp Thr Phe Leu Ala Trp Glu Asn His His Asn Ser Glu Cys Ala Arg Cys Gln Ala Cys Asp Glu Gln Ala Ser Gln Val Ala Leu Glu Asn Cys Ser Ala Val Ala Asp Thr Arg Cys Gly Cys Lys Pro Gly Trp Phe Val Glu Cys Gln Val Ser Gln Cys Val Ser Ser Ser Pro Phe Tyr Cys Gln Pro Cys Leu Asp Cys Gly Ala Leu His Arg His Thr Arg Leu Leu Cys Ser Arg Arg Asp Thr Asp Cys Gly Thr Cys Leu Pro Gly Phe Tyr Glu His Gly Asp Gly Cys Val Ser Cys Pro 

#### Thr

#### (2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 433 base pairs
  - (B) TYPE: Nucleic Acid
  - (C) STRANDEDNESS: Single
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TGCCGGTGAC TTCCACAAGA AGATTGGTCT GTTTTGTTGC AGAGGCTGCC 100
CAGCGGGGCA ACTACCTGAA GGCCCCTTGC ACGGAGCCCT GCGCAACTCC 150
ACCTGCCTTG TGTGTCCCCA AGACACCTTC TTGGCCTGGG AGAACCACCA 200
TAATTCTGAA TGTGCCCGCT GCCAGGCCTG TGATGAGCAG GCCTCCCAGG 250
TGGCGCTGGA GAACTGTTCA GCAGTGGCCG ACACCCGCTG TGGCTGTAAG 300
CAGGGCTGGT TTGTGGAGTG CCAGGGTCAG CCAATGTGTC AGCAGTTTCA 350

# CCCTTCTAAT GCCAACCATG CCTAGACTGC GGGGCCCTGC AACGCAACAC 400 ACGGCTAATN TGTTTCCCGC AGAGATNATT GTT 433

- (2) INFORMATION FOR SEQ ID NO:3:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 28 base pairs
    - (B) TYPE: Nucleic Acid
    - (C) STRANDEDNESS: Single
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

#### CCCGCTGCCA GGCCTGTGAT GAGCAGGC 28

- (2) INFORMATION FOR SEQ ID NO:4:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 28 base pairs
    - (B) TYPE: Nucleic Acid
    - (C) STRANDEDNESS: Single
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

#### CAGGGCCCCG CAGTCTAGGC ATGGTTGG 28

- (2) INFORMATION FOR SEQ ID NO:5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 1438 base pairs
      - (B) TYPE: Nucleic Acid
      - (C) STRANDEDNESS: Single
      - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

GAATTCCGGC GCGGAGGCCG AGAGAGAAGT CACTTGCCCT GGCTCTACCT 50
TGAAGTGGTT CTCAGGGTTG GGGCGAGAGT CGGGGTGGGG ACCGAGATGC 100
AGCTCTATCC TGTGCCCCTG GTCGCAGCAG GCAGCCCAGC GCTTCGCGTG 150
TTCTACTTGG CCTGTCCGCT GCCGCCTAAT GAGCTCAGGT CTAGGCCGAG 200
CAGAGGGGGC ACCTGGTCGG ACTCGGTTGG GCTCGGGGGG CCCCGCCTCC 250
CCCCGCCCGC CAGGCGGGCC CTTCTCGACG GCGCGGGGCG GGCCCTGCGG 300
GCGCGGGGCT GAAGGCGGAA CCACGACGGG CAGAGAGCAC GGAGCCGGGA 350
AGCCCCTGGG CGCCCGTCGG AGGGCTATGG AGCAGCGGCC GCGGGCTGC 400
GCGGCGGTGG CGGCGGCCT CCTCCTGGTG CTGCTGGGG CCCGGGCCCA 450
AGATTGGTCT GTTTTGTTGC AGAGGCTGCC CAGCGGGCA CTACCTGAAG 550

GCCCCTTGCA CGGAGCCCTG CGGCAACTCC ACCTGCCTTG TGTGTCCCCA 600 AGACACCTTC TTGGCCTGGG AGAACCACCA TAATTCTGAA TGTGCCCGCT 650 GCCAGGCCTG TGATGAGCAG GCCTCCCAGG TGGCGCTGGA GAACTGTTCA 700 GCAGTGGCCG ACACCCGCTG TGGCTGTAAG CCAGGCTGGT TTGTGGAGTG 750 CCAGGTCAGC CAATGTGTCA GCAGTTCACC CTTCTACTGC CAACCATGCC 800 TAGACTGCGG GGCCCTGCAC CGCCACACAC GGCTACTCTG TTCCCGCAGA 850 GATACTGACT GTGGGACCTG CCTGCCTGGC TTCTATGAAC ATGGCGATGG 900 CTGCGTGTCC TGCCCCACGT AATTCCTAGC TGTCGTGGGA TGGAGGGAAG 950 GGCGGCTGGG AGCAGAGCAG GGGCCTGGGG TGGGGCAGGT GCTGCTGGTT 1000 CAGGAATAGG AAGAGGGGAT AGGGAGGAGG GAGCCTTGGC CCTGTGATGG 1050 GTGGGCCCCA CTTCAGGCAA ACTTAGATGG CAAAAGAGCA ATCTGGATCC 1100 GCCTTAGCCA GATACATAAG GGTATTTGCC TTCACTTTCA GCCAGCATTC 1150 CCCCCAGCGA TCCTAGCCAG ATATTACAGA TGATTTGTCA CTTACACAGA 1200 GAGTCACATT GATATAGCTT TAAAACTTGG GCTGAAGGAG GTTGAGGCTG 1250 CAGTGAGCTA TGATCGTGCC ACTGCACTTC AGCCTGGGCA ACAGAGCGAG 1300 ACCTATTAAA TAAATAAATA AATATTAAAT CTATTAAATA TTAAATATTA 1350 AATCTATTAA ATAAATAAAT ACAAAGGGCT GAGAGTCAGG ACTGTGCTGC 1400 TAGTTCTCTA GGGGATCTTG GGCAAGTGCA GAGAATTC 1438

# (2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 417 amino acids
  - (B) TYPE: Amino Acid
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
- Met Glu Gln Arg Pro Arg Gly Cys Ala Ala Val Ala Ala Ala Leu
  1 5 10 15
- Leu Leu Val Leu Leu Gly Ala Arg Ala Gln Gly Gly Thr Arg Ser 20 25 30
- Pro Arg Cys Asp Cys Ala Gly Asp Phe His Lys Lys Ile Gly Leu
  35 40 45
- Phe Cys Cys Arg Gly Cys Pro Ala Gly His Tyr Leu Lys Ala Pro 50 55 60
- Cys Thr Glu Pro Cys Gly Asn Ser Thr Cys Leu Val Cys Pro Gln
  65 70 75
- Asp Thr Phe Leu Ala Trp Glu Asn His His Asn Ser Glu Cys Ala 80 85 90

Arg	Cys	Gln	Ala	Cys 95	Asp	Glu	Gln	Ala	Ser 100	Gln	Val	Ala	Leu	Glu 105
Asn	Cys	Ser	Ala	Val 110	Ala	Asp	Thr	Arg	Cys 115	Gly	Cys	Lys	Pro	Gly 120
Trp	Phe	Val	Glu	Cys 125	Gln	Val	Ser	Gln	Cys 130	Val	Ser	Ser	Ser	Pro 135
Phe	Tyr	Cys	Gln	Pro 140	Cys	Leu	Asp	Cys	Gly 145	Ala	Leu	His	Arg	His 150
Thr	Arg	Leu	Leu	Cys 155	Ser	Arg	Arg	Asp	Thr 160	Asp	Cys	Gly	Thr	Cys 165
Leu	Pro	Gly	Phe	Tyr 170	Glu	His	Gly	Asp	Gly 175	Cys	Val	Ser	Cys	Pro 180
Thr	Ser	Thr	Leu	Gly 185	Ser	Cys	Pro	Glu	Arg 190	Cys	Ala	Ala	Val	Cys 195
Gly	Trp	Arg	Gln	Met 200	Phe	Trp	Val	Gln	Val 205	Leu	Leu	Ala	Gly	Leu 210
Val	Val	Pro	Leu	Leu 215	Leu	Gly	Ala	Thr	Leu 220	Thr	Tyr	Thr	Tyr	Arg 225
His	Cys	Trp	Pro	His 230	Lys	Pro	Leu	Val	Thr 235	Ala	Asp	Glu	Ala	Gly 240
Met	Glu	Ala	Leu	Thr 245	Pro	Pro	Pro	Ala	Thr 250	His	Leu	Ser	Pro	Leu 255
Asp	Ser	Ala	His	Thr 260	Leu	Leu	Ala	Pro	Pro 265	Asp	Ser	Ser	Glu	Lys 270
Ile	Cys	Thr	Val	Gln 2 <b>7</b> 5	Leu	Val	Gly	Asn	Ser 280	Trp	Thr	Pro	Gly	Tyr 285
Pro	Glu	Thr	Gln	Glu 290	Ala	Leu	Cys	Pro	Gln 295	Val	Thr	Trp	Ser	Trp 300
Asp	Gln	Leu	Pro	Ser 305	Arg	Ala	Leu	Gly	Pro 310	Ala	Ala	Ala	Pro	Thr 315
Leu	Ser	Pro	Glu	Ser 320	Pro	Ala	Gly	Ser	Pro 325	Ala	Met	Met	Leu	Gln 330
Pro	Gly	Pro	Gln	Leu 335	Tyr	Asp	Val	Met	Asp 340	Ala	Val	Pro	Ala	Arg 345
Arg	Trp	Lys	Glu	Phe 350	Val	Arg	Thr	Leu	Gly 355	Leu	Arg	Glu	Ala	Glu 360
Ile	Glu	Ala	Val	Glu 365	Val	Glu	Ile	Gly	Arg 370	Phe	Arg	Asp	Gln	Gln 375
Tyr	Glu	Met	Leu	Lys 380	Arg	Trp	Arg	Gln	Gln 385	Gln	Pro	Ala	Gly	Leu 390
Gly	Ala	Val	Tyr	Ala 395	Ala	Leu	Glu	Arg	Met 400	Gly	Leu	Asp	Gly	Cys 405

# Val Glu Asp Leu Arg Ser Arg Leu Gln Arg Gly Pro

- (2) INFORMATION FOR SEQ ID NO:7:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 27 base pairs
    - (B) TYPE: Nucleic Acid
    - (C) STRANDEDNESS: Single
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

#### GGCGCTCTGG TGGCCCTTGC AGAAGCC 27

- (2) INFORMATION FOR SEQ ID NO:8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 25 base pairs
    - (B) TYPE: Nucleic Acid
    - (C) STRANDEDNESS: Single
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

#### TTCGGCCGAG AAGTTGAGAA ATGTC 25

- (2) INFORMATION FOR SEQ ID NO:9:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 1634 base pairs
    - (B) TYPE: Nucleic Acid
    - (C) STRANDEDNESS: Single
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

CGGGCCCTGC GGGCGCGGG CTGAAGGCGG AACCACGACG GGCAGAGAGC 50

ACGGAGCCGG GAAGCCCCTG GGCGCCCGTC GGAGGGCT ATG GAG 94
Met Glu
1

CTC CTG GTG CTG GGG GCC CGG GCC CAG GGC GGC ACT 172 Leu Leu Val Leu Gly Ala Arg Ala Gln Gly Gly Thr 20 25

CGT AGC CCC AGG TGT GAC TGT GCC GGT GAC TTC CAC AAG 211
Arg Ser Pro Arg Cys Asp Cys Ala Gly Asp Phe His Lys
30 35 40

AAG ATT GGT CTG TTT TGT TGC AGA GGC TGC CCA GCG GGG 250 Lys Ile Gly Leu Phe Cys Cys Arg Gly Cys Pro Ala Gly 45

			TGC Cys					289
			CCC Pro				_	328
			TCT Ser					367
			TCC Ser 100					406
			ACC Thr		_			445
			CAG Gln					484
			CAA Gln					523
			CGG Arg					562
			TGC Cys 165					601
			TCC Ser					640
			TGT Cys					679
			CAG Gln					718
			GGG Gly			_	_	757
			CAC His 230					796
			GCT Ala					835
			GAC Asp					874

TTG GTG GGT AAC AGC TGG ACC CCT GGC TAC CCC GAG ACC 952 Leu Val Gly Asn Ser Trp Thr Pro Gly Tyr Pro Glu Thr 280 285
CAG GAG GCG CTC TGC CCG CAG GTG ACA TGG TCC TGG GAC 991 Gln Glu Ala Leu Cys Pro Gln Val Thr Trp Ser Trp Asp 290 295 300
CAG TTG CCC AGC AGA GCT CTT GGC CCC GCT GCT GCG CCC 1030 Gln Leu Pro Ser Arg Ala Leu Gly Pro Ala Ala Pro 305 310
ACA CTC TCG CCA GAG TCC CCA GCC GGC TCG CCA GCC ATG 1069 Thr Leu Ser Pro Glu Ser Pro Ala Gly Ser Pro Ala Met 315 320 325
ATG CTG CAG CCG GGC CCG CAG CTC TAC GAC GTG ATG GAC 1108  Met Leu Gln Pro Gly Pro Gln Leu Tyr Asp Val Met Asp  330 335 340
GCG GTC CCA GCG CGC CGC TGG AAG GAG TTC GTG CGC ACG 1147 Ala Val Pro Ala Arg Arg Trp Lys Glu Phe Val Arg Thr 345 350
CTG GGG CTG CGC GAG GCA GAG ATC GAA GCC GTG GAG GTG 1186 Leu Gly Leu Arg Glu Ala Glu Ile Glu Ala Val Glu Val 355 360 365
GAG ATC GGC CGC TTC CGA GAC CAG CAG TAC GAG ATG CTC 1225 Glu Ile Gly Arg Phe Arg Asp Gln Gln Tyr Glu Met Leu 370 375
AAG CGC TGG CGC CAG CAG CCC GCG GGC CTC GGA GCC 1264 Lys Arg Trp Arg Gln Gln Gln Pro Ala Gly Leu Gly Ala 380 385 390
GTT TAC GCG GCC CTG GAG CGC ATG GGG CTG GAC GGC TGC 1303 Val Tyr Ala Ala Leu Glu Arg Met Gly Leu Asp Gly Cys 395 400 405
GTG GAA GAC TTG CGC AGC CGC CTG CAG CGC GGC CCG T 1340 Val Glu Asp Leu Arg Ser Arg Leu Gln Arg Gly Pro 410 415 417
GACACGGCGC CCACTTGCCA CCTAGGCGCT CTGGTGGCCC TTGCAGAAGC 1390
CCTAAGTACG GTTACTTATG CGTGTAGACA TTTTATGTCA CTTATTAAGC 1440
CGCTGGCACG GCCCTGCGTA GCAGCACCAG CCGGCCCCAC CCCTGCTCGC 1490
CCCTATCGCT CCAGCCAAGG CGAAGAAGCA CGAACGAATG TCGAGAGGGG 1540 GTGAAGACAT TTCTCAACTT CTCGGCCGGA GTTTGGCTGA GATCGCGGTA 1590
TTAAATCTGT GAAAGAAAAC AAAAAAAAAAA AAAAAAAAA AAAA

# (2) INFORMATION FOR SEQ ID NO:10:

# (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 29 base pairs
- (B) TYPE: Nucleic Acid
- (C) STRANDEDNESS: Single
- (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

#### ATCAGGGACT TTCCGCTGGG GACTTTCCG 29

- (2) INFORMATION FOR SEQ ID NO:11:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 30 base pairs
    - (B) TYPE: Nucleic Acid
    - (C) STRANDEDNESS: Single
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

#### AGGATGGGAA GTGTGTGATA TATCCTTGAT 30

- (2) INFORMATION FOR SEQ ID NO:12:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 154 amino acids
    - (B) TYPE: Amino Acid
    - (D) TOPOLOGY: Linear
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:
- Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile
  1 5 10 15
- Cys Cys Thr Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys
  20 25 30
- Pro Gly Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly 35 40 45
- Ser Phe Thr Ala Ser Glu Asn His Leu Arg His Cys Leu Ser Cys
  50 55 60
- Ser Lys Cys Arg Lys Glu Met Gly Gln Val Glu Ile Ser Ser Cys
  65 70 75
- Thr Val Asp Arg Asp Thr Val Cys Gly Cys Arg Lys Asn Gln Tyr 80 85 90
- Arg His Tyr Trp Ser Glu Asn Leu Phe Gln Cys Phe Asn Cys Ser 95 100 105
- Leu Cys Leu Asn Gly Thr Val His Leu Ser Cys Gln Glu Lys Gln
  110 115 120
- Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu Asn 125 130 135
- Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys Thr 140 145 150

Lys Leu Cys Leu

#### (2) INFORMATION FOR SEQ ID NO:13:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 163 amino acids
  - (B) TYPE: Amino Acid
  - (D) TOPOLOGY: Linear

### (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Thr Cys Arg Leu Arg Glu Tyr Tyr Asp Gln Thr Ala Gln Met Cys
1 5 10 15

Cys Ser Lys Cys Ser Pro Gly Gln His Ala Lys Val Phe Cys Thr 20 25 30

Lys Thr Ser Asp Thr Val Cys Asp Ser Cys Glu Asp Ser Thr Tyr 35 40 45

Thr Gln Leu Trp Asn Trp Val Pro Glu Cys Leu Ser Cys Gly Ser 50 60

Arg Cys Ser Ser Asp Gln Val Glu Thr Gln Ala Cys Thr Arg Glu
65 70 75

Gln Asn Arg Ile Cys Thr Cys Arg Pro Gly Trp Tyr Cys Ala Leu 80 85 90

Ser Lys Gln Glu Gly Cys Arg Leu Cys Ala Pro Leu Arg Lys Cys 95 100 105

Arg Pro Gly Phe Gly Val Ala Arg Pro Gly Thr Glu Thr Ser Asp 110 115 120

Val Val Cys Lys Pro Cys Ala Pro Gly Thr Phe Ser Asn Thr Thr
125 130 135

Ser Ser Thr Asp Ile Cys Arg Pro His Gln Ile Cys Asn Val Val 140 145 150

Ala Ile Pro Gly Asn Ala Ser Arg Asp Ala Val Cys Thr 155 160

#### (2) INFORMATION FOR SEQ ID NO:14:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 170 amino acids
  - (B) TYPE: Amino Acid
  - (D) TOPOLOGY: Linear

# (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Thr Cys Arg Asp Gln Glu Lys Glu Tyr Tyr Glu Pro Gln His Arg
1 5 10 15

Ile Cys Cys Ser Arg Cys Pro Pro Gly Thr Tyr Val Ser Ala Lys
20 25 30

Cys Ser Arg Ile Arg Asp Thr Val Cys Ala Thr Cys Ala Glu Asn
35 40 45

Ser Tyr Asn Glu His Trp Asn Tyr Leu Thr Ile Cys Gln Leu Cys
50 55 60

Arg Pro Cys Asp Pro Val Met Gly Leu Glu Glu Ile Ala Pro Cys Thr Ser Lys Arg Lys Thr Gln Cys Arg Cys Gln Pro Gly Met Phe Cys Ala Ala Trp Ala Leu Glu Cys Thr His Cys Glu Leu Leu Ser Asp Cys Pro Pro Gly Thr Glu Ala Glu Leu Lys Asp Glu Val Gly Lys Gly Asn Asn His Cys Val Pro Cys Lys Ala Gly His Phe Gln Asn Thr Ser Ser Pro Ser Ala Arg Cys Gln Pro His Thr Arg Cys Glu Asn Gln Gly Leu Val Glu Ala Ala Pro Gly Thr Ala Gln Ser Asp Thr Thr Cys Lys 

#### (2) INFORMATION FOR SEQ ID NO:15:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 119 amino acids
  - (B) TYPE: Amino Acid
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Asn Leu Glu Gly Leu His His Asp Gly Gln Phe Cys His Lys Pro 

Cys Pro Pro Gly Glu Arg Lys Ala Arg Asp Cys Thr Val Asn Gly 

Asp Glu Pro Asp Cys Val Pro Cys Gln Glu Gly Lys Glu Tyr Thr 

Asp Lys Ala His Phe Ser Ser Lys Cys Arg Arg Cys Arg Leu Cys 

Asp Glu Gly His Gly Leu Glu Val Glu Ile Asn Cys Thr Arg Thr 

Gln Asn Thr Lys Cys Arg Cys Lys Pro Asn Phe Phe Cys Asn Ser 

Thr Val Cys Glu His Cys Asp Pro Cys Thr Lys Cys Glu His Gly 

Ile Ile Lys Glu Cys Thr Leu Thr Ser Asn Thr Lys Cys Lys 

#### (2) INFORMATION FOR SEQ ID NO:16:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 159 amino acids
  - (B) TYPE: Amino Acid
  - (D) TOPOLOGY: Linear

# (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Ala Cys Pro Thr Gly Leu Tyr Thr His Ser Gly Glu Cys Cys Lys Ala Cys Asn Leu Gly Glu Gly Val Ala Gln Pro Cys Gly Ala Asn Gln Thr Val Cys Glu Pro Cys Leu Asp Ser Val Thr Phe Ser Asp Val Val Ser Ala Thr Glu Pro Cys Lys Pro Cys Thr Glu Cys Val Gly Leu Gln Ser Met Ser Ala Pro Cys Val Glu Ala Asp Asp Ala Val Cys Arg Cys Ala Tyr Gly Tyr Tyr Gln Asp Glu Thr Thr Gly Arg Cys Glu Ala Cys Arg Val Cys Glu Ala Gly Ser Gly Leu Val Phe Ser Cys Gln Asp Lys Gln Asn Thr Val Cys Glu Glu Cys Pro Asp Gly Thr Tyr Ser Asp Glu Ala Asn His Val Asp Pro Cys Leu Pro Cys Thr Val Cys Glu Asp Thr Glu Arg Gln Leu Arg Glu Cys 

# (2) INFORMATION FOR SEQ ID NO:17:

(i) SEQUENCE CHARACTERISTICS:

Thr Arg Trp Ala Asp Ala Glu Cys Glu 

- (A) LENGTH: 163 amino acids
- (B) TYPE: Amino Acid
- (D) TOPOLOGY: Linear

#### (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Ala Cys Arg Glu Lys Gln Tyr Leu Ile Asn Ser Gln Cys Cys Ser Leu Cys Gln Pro Gly Gln Lys Leu Val Ser Asp Cys Thr Glu Phe Thr Glu Thr Glu Cys Leu Pro Cys Gly Glu Ser Glu Phe Leu Asp Thr Trp Asn Arg Glu Thr His Cys His Gln His Lys Tyr Cys Asp Pro Asn Leu Gly Leu Arg Val Gln Gln Lys Gly Thr Ser Glu Thr Asp Thr Ile Cys Thr Cys Glu Glu Gly Trp His Cys Thr Ser Glu 

Ala Cys Glu Ser Cys Val Leu His Arg Ser Cys Ser Pro Gly Phe 105 95 100 Gly Val Lys Gln Ile Ala Thr Gly Val Ser Asp Thr Ile Cys Glu 120 110 115 Pro Cys Pro Val Gly Phe Phe Ser Asn Val Ser Ser Ala Phe Glu 125 130 135 Lys Cys His Pro Trp Thr Ser Cys Glu Thr Lys Asp Leu Val Val 140 150 145 Gln Gln Ala Gly Thr Asn Lys Thr Asp Val Val Cys Gly

160

#### (2) INFORMATION FOR SEQ ID NO:18:

(i) SEQUENCE CHARACTERISTICS:

155

- (A) LENGTH: 121 amino acids
- (B) TYPE: Amino Acid
- (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Ser Cys Pro Glu Arg His Tyr Trp Ala Gln Gly Lys Leu Cys Cys 15

Gln Met Cys Glu Pro Gly Thr Phe Leu Val Lys Asp Cys Asp Gln 30

His Arg Lys Ala Ala Gln Cys Asp Pro Cys 10

Phe Ser Pro Asp His His Thr Arg Pro His Cys Glu Ser Cys Arg

50 Fig. 55 Fig. 55 Fig. 56 Fig. 57 Fig. 58 Fig. 60

His Cys Asn Ser Gly Leu Leu Val Arg Asn Cys Thr Ile Thr Ala
65 70 75

Asn Ala Glu Cys Ala Cys Arg Asn Gly Trp Gln Cys Arg Asp Lys 80 85 90

Glu Cys Thr Glu Cys Asp Pro Leu Pro Asn Pro Ser Leu Thr Ala 95 100 105

Arg Ser Ser Gln Ala Leu Ser Pro His Pro Gln Pro Thr His Leu 110 115 120

Pro

# (2) INFORMATION FOR SEQ ID NO:19:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 123 amino acids
  - (B) TYPE: Amino Acid
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Thr Cys His Gly Asn Pro Ser His Tyr Tyr Asp Lys Ala Val Arg
1 1 10 15

Arg Cys Cys Tyr Arg Cys Pro Met Gly Leu Phe Pro Thr Gln Gln Cys Pro Gln Arg Pro Thr Asp Cys Arg Lys Gln Cys Glu Pro Asp Tyr Tyr Leu Asp Glu Ala Asp Arg Cys Thr Ala Cys Val Thr Cys Ser Arg Asp Asp Leu Val Glu Lys Thr Pro Cys Ala Trp Asn Ser Ser Arg Val Cys Glu Cys Arg Pro Gly Met Phe Cys Ser Thr Ser Ala Val Asn Ser Cys Ala Arg Cys Phe Phe His Ser Val Cys Pro Ala Gly Met Ile Val Lys Phe Pro Gly Thr Ala Gln Lys Asn Thr Val Cys Glu

#### (2) INFORMATION FOR SEQ ID NO:20:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 138 amino acids
  - (B) TYPE: Amino Acid
  - (D) TOPOLOGY: Linear

Ile Cys Glu

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:
- His Cys Val Gly Asp Thr Tyr Pro Ser Asn Asp Arg Cys Cys His Glu Cys Arg Pro Gly Asn Gly Met Val Ser Arg Cys Ser Arg Ser Gln Asn Thr Val Cys Arg Pro Cys Gly Pro Gly Phe Tyr Asn Asp Val Val Ser Ser Lys Pro Cys Lys Pro Cys Thr Trp Cys Asn Leu Arg Ser Gly Ser Glu Arg Lys Gln Leu Cys Thr Ala Thr Gln Asp Thr Val Cys Arg Cys Arg Ala Gly Thr Gln Pro Leu Asp Ser Tyr Lys Pro Gly Val Asp Cys Ala Pro Cys Pro Pro Gly His Phe Ser Pro Gly Asp Asn Gln Ala Cys Lys Pro Trp Thr Asn Cys Thr Leu Ala Gly Lys His Thr Leu Gln Pro Ala Ser Asn Ser Ser Asp Ala

#### (2) INFORMATION FOR SEQ ID NO:21:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 197 amino acids
  - (B) TYPE: PRT
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Met Gly Leu Ser Thr Val Pro Asp Leu Leu Leu Pro Leu Val Leu
1 5 10 15

Leu Glu Leu Val Gly Ile Tyr Pro Ser Gly Val Ile Gly Leu
20 25 30

Val Pro His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys Pro 35 40 45

Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr
50 55 60

Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro 65 70 75

Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr 80 85 90

Ala Ser Glu Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys
95 100 105

Arg Lys Glu Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp 110 115 120

Arg Asp Thr Val Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr
125 130 135

Trp Ser Glu Asn Leu Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu 140 145 150

Asn Gly Thr Val His Leu Ser Cys Gln Glu Lys Gln Asn Thr Val 155 160 165

Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu Asn Glu Cys Val 170 175 180

Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys Thr Lys Leu Cys 185 190 195

Leu Pro

#### (2) INFORMATION FOR SEQ ID NO:22:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 167 amino acids
  - (B) TYPE: PRT
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

Met Leu Gly Ile Trp Thr Leu Leu Pro Leu Val Leu Thr Ser Val
1 5 10 15

Ala Arg Leu Ser Ser Lys Ser Val Asn Ala Gln Val Thr Asp Ile 30 20 Asn Ser Lys Gly Leu Glu Leu Arg Lys Thr Val Thr Thr Val Glu 45 35 40 Thr Gln Asn Leu Glu Gly Leu His His Asp Gly Gln Phe Cys His 50 60 55 Lys Pro Cys Pro Pro Gly Glu Arg Lys Ala Arg Asp Cys Thr Val 75 65 Asn Gly Asp Glu Pro Asp Cys Val Pro Cys Gln Glu Gly Lys Glu 80 85 90 Tyr Thr Asp Lys Ala His Phe Ser Ser Lys Cys Arg Arg Cys Arg 95 100 105 Leu Cys Asp Glu Gly His Gly Leu Glu Val Glu Ile Asn Cys Thr 110 115 120 Arg Thr Gln Asn Thr Lys Cys Arg Cys Lys Pro Asn Phe Phe Cys 125 130 135 Asn Ser Thr Val Cys Glu His Cys Asp Pro Cys Thr Lys Cys Glu 150 140 145 His Gly Ile Ile Lys Glu Cys Thr Leu Thr Ser Asn Thr Lys Cys 155 160 165 Lys Glu

#### (2) INFORMATION FOR SEQ ID NO:23:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 78 amino acids
  - (B) TYPE: PRT
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

Val Val Glu Asn Val Pro Pro Leu Arg Trp Lys Glu Phe Val Arg
1 10 15

Arg Leu Gly Leu Ser Asp His Glu Ile Asp Arg Leu Glu Leu Gln
20 25 30

Asn Gly Arg Cys Leu Arg Glu Ala Gln Tyr Ser Met Leu Ala Thr

Trp Arg Arg Thr Pro Arg Arg Glu Ala Thr Leu Glu Leu Leu 50 55 60

Gly Arg Val Leu Arg Asp Met Asp Leu Leu Gly Cys Leu Glu Asp
65 70 75

Ile Glu Glu

# (2) INFORMATION FOR SEQ ID NO:24:

#### (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 77 amino acids
- (B) TYPE: PRT
- (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Ile Ala Gly Val Met Thr Leu Ser Gln Val Lys Gly Phe Val Arg
1 5 10 15

Lys Asn Gly Val Asn Glu Ala Lys Ile Asp Glu Ile Lys Asn Asp

Asn Val Gln Asp Thr Ala Glu Gln Lys Val Gln Leu Leu Arg Asn
35 40 45

Trp His Gln Leu His Gly Lys Lys Glu Ala Tyr Asp Thr Leu Ile
50 55 60

Lys Asp Leu Lys Lys Ala Asn Leu Cys Thr Leu Ala Glu Lys Ile
65 70 75

Gln Thr

# (2) INFORMATION FOR SEQ ID NO:25:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 74 amino acids
  - (B) TYPE: PRT
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

Ile Cys Asp Asn Val Gly Lys Asp Trp Arg Arg Leu Ala Arg Gln
1 5 10 15

Leu Lys Val Ser Asp Thr Lys Ile Asp Ser Ile Glu Asp Arg Tyr
20 25 30

Pro Arg Asn Leu Thr Glu Arg Val Arg Glu Ser Leu Arg Ile Trp
35 40 45

Lys Asn Thr Glu Lys Glu Asn Ala Thr Val Ala His Leu Val Gly
50 55 60

Ala Leu Arg Ser Cys Gln Met Asn Leu Val Ala Asp Leu Val
65 70

#### (2) INFORMATION FOR SEQ ID NO:26:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 77 amino acids
  - (B) TYPE: PRT
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Asn Arg Pro Leu Ser Leu Lys Asp Gln Gln Thr Phe Ala Arg Ser
1 5 10 15

Val Gly Leu Lys Trp Arg Lys Val Gly Arg Ser Leu Gln Arg Gly
20 25 30

Cys Arg Ala Leu Arg Asp Pro Ala Leu Asp Ser Leu Ala Tyr Glu
35 40 45

Tyr Glu Arg Glu Gly Leu Tyr Glu Gln Ala Phe Gln Leu Leu Arg
50 55 60

Arg Phe Val Gln Ala Glu Gly Arg Arg Ala Thr Leu Gln Arg Leu 65 70 75

Val Glu

#### (2) INFORMATION FOR SEQ ID NO:27:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 77 amino acids
  - (B) TYPE: PRT
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

Ile Arg Glu Asn Leu Gly Lys His Trp Lys Asn Cys Ala Arg Lys
1 5 10 15

Leu Gly Phe Thr Gln Ser Gln Ile Asp Glu Ile Asp His Asp Tyr
20 25 30

Glu Arg Asp Gly Leu Lys Glu Lys Val Tyr Gln Met Leu Gln Lys
35 40 45

Trp Val Met Arg Glu Gly Ile Lys Gly Ala Thr Val Gly Lys Leu
50 55 60

Ala Gln Ala Leu His Gln Cys Ser Arg Ile Asp Leu Leu Ser Ser 75

Leu Thr

# (2) INFORMATION FOR SEQ ID NO:28:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 63 amino acids
  - (B) TYPE: PRT
  - (D) TOPOLOGY: Linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

Met Ala Val Ala Phe Tyr Ile Pro Asp Gln Ala Thr Leu Leu Arg
1 5 10 15

Glu Ala Glu Gln Lys Glu Gln Gln Ile Leu Arg Leu Arg Glu Ser 20 25 30

Gln Trp Arg Phe Leu Ala Thr Val Val Leu Glu Thr Leu Lys Gln
35 40 45

Tyr Thr Ser Cys His Pro Lys Thr Gly Arg Lys Ser Gly Lys Tyr
50 55 60

Arg Lys Pro